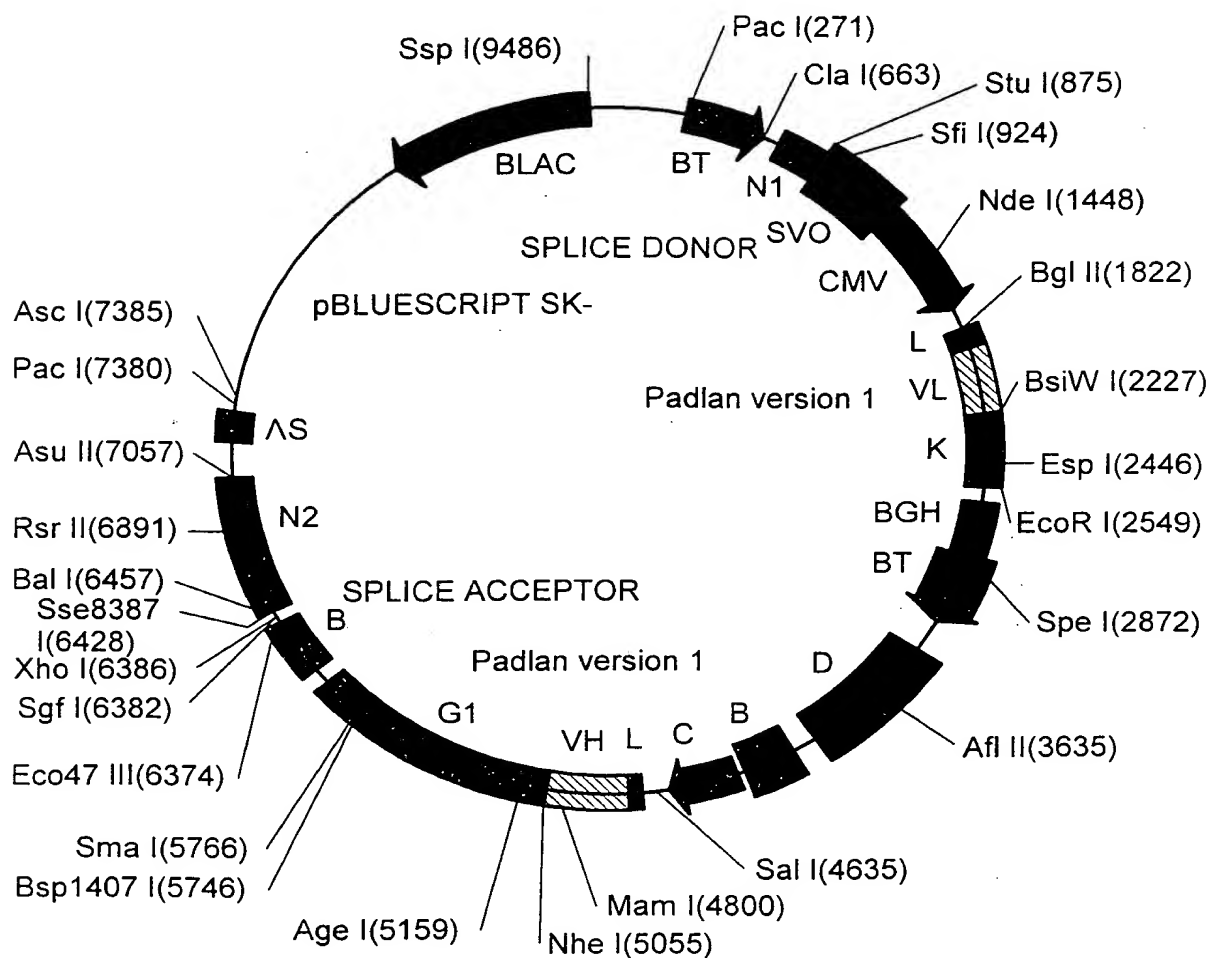


FIG. 1

Anti-GP39 HU24-31 in N5KG1 (Version #1)



9597 BP

Map BY Mitchell Raff

06/01/95

C = Cytomegalovirus promoter/enhancer

BT = Mouse Beta globin major promoter

B = Bovine growth hormone polyadenylation

N1 = Neomycin phosphotransferase exon 1

N2 = Neomycin phosphotransferase exon 2

K = Human immunoglobulin kappa constant region

G1 = Human immunoglobulin gamma 1 constant region

VL = Anti-GP39 variable light region (version #1)

VH = Anti-GP39 variable heavy region (version #1)

NONCUTTERS = AvrII, BstI 107I, DraIII, FseI, HindIII, I-PpoI, I-SceI,

KpnI, MluI, MunI, PmeI, PmlI, SgrAI, SrfI, Swa I, XbaI, XcmI

N5KG1 cut BglII + BsiWI and VL dropped in & cut SalI + NheI and VH dropped in.

L = Leader

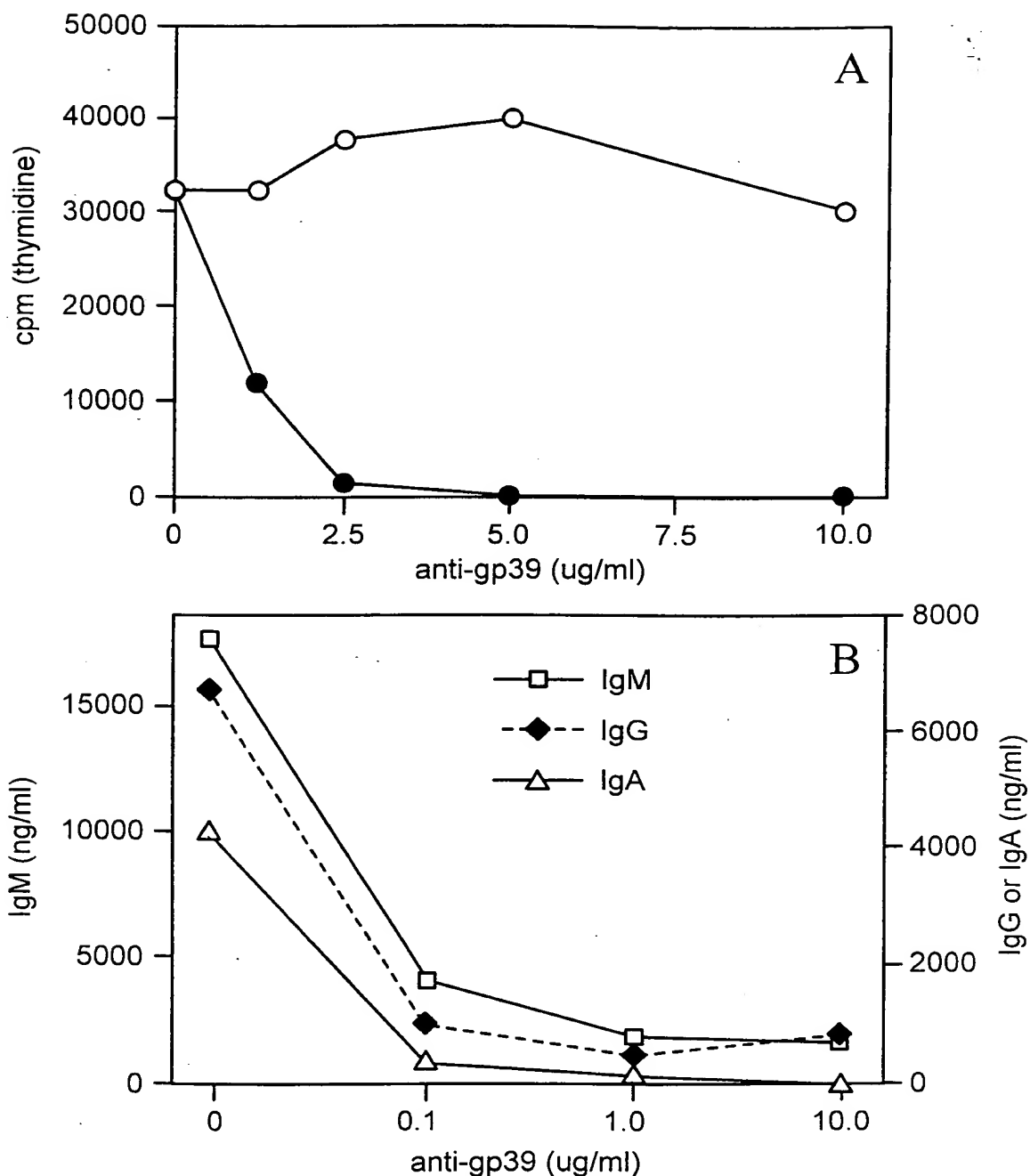
SVO = SV40 origin

SV = SV40 polyadenylation

D = Dihydrofolate Reductase

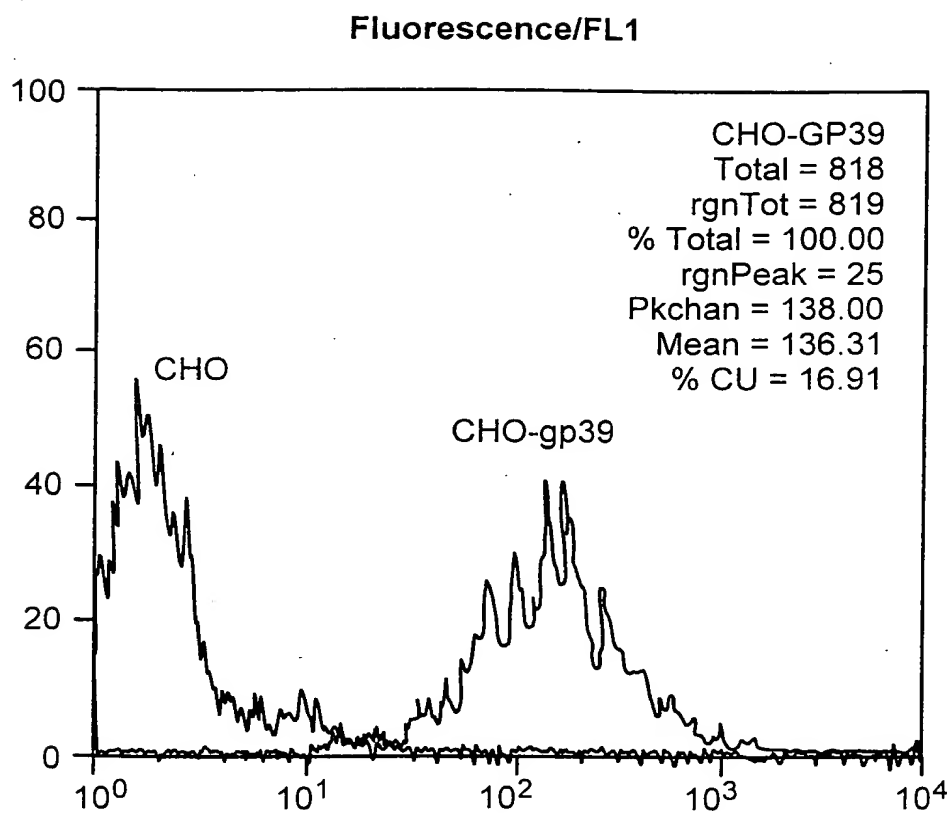
BLAC = Betalactamase gene

FIG. 2



Anti-gp39 inhibits B cell proliferation and differentiation, but not allogeneic T cell proliferation. A. Human PBL were cultured in 96 well plates (0.1×10^6 /well) in the presence or absence of the 20% (v/v) soluble gp39-CD8 (sgp39-CD8) fusion protein and 5 ug/ml rhIL-4 for 3 d. Anti-gp39 mAb, 24-31 (●), or a control murine IgG1 mAb (○), were added at a range of concentrations (1.25-10 ug/ml). Cultures were pulsed with 1 μ Ci 3 H-thymidine during the final 6 hr of a 72 hr culture period. B. Mitomycin treated T cells (5×10^4 /well) activated with immobilized anti-CD3 (64.1) were cultured with 2.5×10^4 /well IgD⁺ B cells in 96-well microtiter plates for 12d in the presence or absence of various concentrations (0.1-10.0 ug/ml) of anti-gp39 mAb, 24-31. Culture supernatants were subsequently assayed for IgM (□), IgG (◆), and IgA (△) by isotype specific ELISA.

FIG. 3



FACS analysis of non-transfected CHO cells and a gp39 transfectant. 1×10^6 cells were treated with the mouse anti-gp39 antibody 24-31 and then with a goat-anti-mouse IgG-FITC conjugate (Southern Biotechnology Associates). The samples were analyzed on FACScan (Becton Dickenson).

0987441-060601

24-31 Humanized V_L #1

| | | | | | | | | | | | | | |
|--------|-----|-----|-----|------|-----|------|-----|-------|-----|------|-----|-----|-----|
| BglIII | | 9 | | 18 | | 27 | | 36 | | 45 | | 54 | |
| AGA | TCT | CTC | ACC | ATG | GGC | TTC | AAG | ATG | GAG | TCA | CAG | TTT | GTA |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| | | | | M | G | F | K | M | E | S | Q | F | V |
| 63 | | 72 | | 81 | | FR1 | | 90 | | 99 | | 108 | |
| GCG | TTT | CTC | TGG | TTG | TCT | GGT | GTT | GAT | GGA | GAC | ATT | GTG | ATG |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| A | F | L | W | L | S | G | V | D | G | D | I | V | M |
| 117 | | 126 | | 135 | | 144 | | 153 | | CDR1 | | 162 | |
| TCT | TTC | CTC | TCC | GCC | TCC | GTA | GGA | GAC | AGG | GTC | ACC | ATC | ACC |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| S | F | L | S | A | S | V | G | D | R | V | T | I | T |
| 171 | | 180 | | 189 | | FR2 | | 198 | | 207 | | 216 | |
| CAG | AAT | GTG | ATT | ACT | GCT | GTA | GCC | TGG | TAT | CAA | CAG | AAA | CCA |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Q | N | V | I | T | A | V | A | W | Y | Q | Q | K | P |
| 225 | | 234 | | CDR2 | | 243 | | 252 | | FR3 | | 261 | |
| AAA | TTG | CTG | ATT | TAC | TCG | GCA | TCC | AAT | CGG | TAC | ACT | GGA | GTC |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| K | L | L | I | Y | S | A | S | N | R | Y | T | G | V |
| 279 | | 288 | | 297 | | 306 | | 315 | | 324 | | 333 | |
| TCA | GGC | AGT | GGG | TCT | GGG | ACA | GAT | TTC | ACT | CTC | ACC | ATC | AGC |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| S | G | S | G | S | G | T | D | F | T | L | T | I | S |
| 333 | | 342 | | 351 | | CDR3 | | 360 | | 369 | | 378 | |
| GAA | GAC | TTC | GCA | GAT | TAT | TTC | TGC | CAG | CAA | TAT | AAC | AGC | TAT |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| E | D | F | A | D | Y | F | C | Q | Q | Y | N | S | Y |
| FR4 | | 387 | | 396 | | 405 | | BsiWI | | 3 | | 378 | |
| GGA | GGG | GGG | ACC | AAG | CTG | GAA | ATC | AAA | CGT | ACG | --- | --- | --- |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| G | G | G | T | K | L | E | I | K | R | T | | | |

FIGURE 4

24-31 Humanized V_L #2

| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
|----|---------------|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|-----|-----|-----|-----|--------------|-----|-----|-----|-----|-----|-----|--|--|-----|------|--|--|--|-----|-----|
| 5' | <u>BglIII</u> | | | | 9 | | | | | 18 | | | | | 27 | | | | | 36 | | | | | 45 | | | | | 54 | |
| | AGA | TCT | CTC | ACC | | ATG | GGC | TTC | AAG | ATG | GAG | TCA | CAG | TTT | CTG | GCC | TTT | GTA | TTC | | | | | | | | | | | | |
| | | | | | | M | G | F | K | M | E | S | Q | F | L | A | F | V | F | | | | | | | | | | | | |
| | | | | | 63 | | | | | 72 | | | | | 81 | | | | | 90 | FR1 | | | | | 99 | | | | | 108 |
| | GCG | TTT | CTC | TGG | TTG | TCT | GGT | GTT | GAT | GGA | GAC | ATT | GTG | ATG | ACC | CAG | TCT | CCA | | | | | | | | | | | | | |
| | A | F | L | W | L | S | G | V | D | G | D | I | V | M | T | Q | S | P | | | | | | | | | | | | | |
| | | | | | 117 | | | | | 126 | | | | | 135 | | | | | 144 | | | | | 153 | CDR1 | | | | 162 | |
| | GAT | TCT | CTC | GCC | GTG | TCC | CTC | GGA | GAG | AGG | GCC | ACC | ATC | AAC | TGC | AAG | GCC | AGT | | | | | | | | | | | | | |
| | D | S | L | A | V | S | L | G | E | R | A | T | I | N | C | K | A | S | | | | | | | | | | | | | |
| | | | | | 171 | | | | | 180 | | | | | 189 | FR2 | | | | | 198 | | | | | 207 | | | | | 216 |
| | CAG | AAT | GTG | ATT | ACT | GCT | GTA | GCC | TGG | TAT | CAA | CAG | AAA | CCA | GGA | CAA | TCT | CCT | | | | | | | | | | | | | |
| | Q | N | V | I | T | A | V | A | W | Y | Q | Q | K | P | G | Q | S | P | | | | | | | | | | | | | |
| | | | | | 225 | | | | | 234 | CDR2 | | | | | 243 | | | | | 252 | FR3 | | | | 261 | | | | | 270 |
| | AAA | TTG | CTG | ATT | TAC | TCG | GCA | TCC | AAT | CGG | TAC | ACT | GGA | GTC | CCT | GAT | CGC | TTC | | | | | | | | | | | | | |
| | K | L | L | I | Y | S | A | S | N | R | Y | T | G | V | P | D | R | F | | | | | | | | | | | | | |
| | | | | | 279 | | | | | 288 | | | | | 297 | | | | | 306 | | | | | 315 | | | | | 324 | |
| | TCA | GGC | AGT | GGG | TCT | GGG | ACA | GAT | TTC | ACT | CTC | ACC | ATC | AGC | TCT | CTC | CAG | GCC | | | | | | | | | | | | | |
| | S | G | S | G | S | G | T | D | F | T | L | T | I | S | S | L | Q | A | | | | | | | | | | | | | |
| | | | | | 333 | | | | | 342 | | | | | 351 | CDR3 | | | | | 360 | | | | | 369 | | | | | 378 |
| | GAA | GAC | GTG | GCA | GAT | TAT | TTC | TGC | CAG | CAA | TAT | AAC | AGC | TAT | CCG | TAC | ACG | TTC | | | | | | | | | | | | | |
| | E | D | V | A | D | Y | F | C | Q | Q | Y | N | S | Y | P | Y | T | F | | | | | | | | | | | | | |
| | FR4 | | | | 387 | | | | | 396 | | | | | 405 | <u>BsiWI</u> | | | | | | | | | | | | | | | |
| | GGA | GGG | GGG | ACC | AAG | CTG | GAA | ATC | AAA | CGT | ACG | 3' | | | | | | | | | | | | | | | | | | | |
| | G | G | G | T | K | L | E | I | K | R | T | | | | | | | | | | | | | | | | | | | | |

FIGURE 5

24-31 Humanized V_H #1

| | | | | | | | | | | | | | | |
|-------------|---------|-------------|-------------|-------------|-------------|-------------|------|------|--|----|--|----|--|--|
| SalI | | 9 | | 18 | | 27 | | 36 | | 45 | | 54 | | |
| 5' | GTC GAC | ATG ATG GTG | TTA AGT CTT | CTG TAC CTG | TTG ACA GCC | CTT CCG GGT | TTC | | | | | | | |
| | | M M V | L S L | L Y L | L T A | L P G | F | | | | | | | |
| CTG TCA | | 63 FR1 | 72 | 81 | 90 | 99 | 108 | | | | | | | |
| GAG GTG CAG | | CTT CAG GAG | TCA GGA CCT | GGC CTC GTG | AAA CCT TCT | GAG | | | | | | | | |
| L S | | E V Q | L Q E | S G P | G L V | K P S | E | | | | | | | |
| ACT CTG | | 117 | 126 | 135 | 144 | 153 CDR1 | 162 | | | | | | | |
| TCC CTC ACC | | TGT ACC GTC | TCT GGC GAC | TCC ATC ACT | AAT GGT TTC | TGG | | | | | | | | |
| T L S | | L T C | T V S | G D S | I T | N G F | W | | | | | | | |
| ATC TGG | | 171 FR2 | 180 | 189 | 198 | 207 | CDR2 | 216 | | | | | | |
| ATC CGG AAA | | CCA CCA GGG | AAT AAA CTT | GAG TAC | ATG GGC | TAC ATA AGT | | | | | | | | |
| I W | | I R K | P P G | N K L | E Y M | G Y I | S | | | | | | | |
| TAC AGT | | 225 | 234 | 243 | 252 | 261 FR3 | 270 | | | | | | | |
| GGT AGC ACT | | TAC TAC AAT | CCA TCT CTC | AAG AGT | CGA ATC TCC | ATC TCT | | | | | | | | |
| Y S G | | S T Y | Y N P | S L K | S R | I S I | S | | | | | | | |
| CGC GAC | | 279 | 288 | 297 | 306 | 315 | 324 | | | | | | | |
| ACA TCC AAG | | AAC CAG TTC | TCT CTA AAG | TTG TCT TCT | GTG ACT GCC | GCC | | | | | | | | |
| R D T | | S K N | Q F S | L K L | S S V | T A A | | | | | | | | |
| GAC ACA | | 333 | 342 | 351 | CDR3 | 360 | 369 | 378 | | | | | | |
| GGC GTG TAT | | TAC TGT GCC | TGC CGC AGT | TAC GGG AGG | ACC CCG TAC | TAC | | | | | | | | |
| D T G | | V Y Y | C A C | R S Y | G R T | P Y Y | | | | | | | | |
| TTT GAC | | 387 | FR4 | 396 | 405 | 414 | 423 | NheI | | | | | | |
| TTC TGG GGC | | CAA GGC ACC | ACT CTC ACC | GTC TCC TCA | GCT AGC | 3' | | | | | | | | |
| F D F | | W G Q | G T T | L T V | S S A | S | | | | | | | | |

FIGURE 6

Anti-gp39 24-31 V_K Sequence

| | | | | | | | | | | | | | | | | | | | | | |
|----|--------|-----|-----|-----|-----|-----|------|-----|-----|------|-----|-----|--------|-----|-----|------|-----|-----|-----|--|--|
| 5' | Bgl II | | | 9 | | | 18 | | | 27 | | | 36 | | | 45 | | | 54 | | |
| | AGA | TCT | CTC | ACC | ATG | GGC | TTC | AAG | ATG | GAG | TCA | CAG | TTT | CTG | GCC | TTT | GTA | TTC | | | |
| | | | | | M | G | F | K | M | E | S | Q | F | L | A | F | V | F | | | |
| | 63 | | | 72 | | | 81 | | | +1 | | | 90 | | | 99 | | | 108 | | |
| | GCG | TTT | CTC | TGG | TTG | TCT | GGT | GTT | GAT | GGA | GAC | ATT | GTG | ATG | ACC | CAG | TCT | CAA | | | |
| | A | F | L | W | L | S | G | V | D | G | D | I | V | M | T | Q | S | Q | | | |
| | 117 | | | 126 | | | 135 | | | 144 | | | 153 | | | CDR1 | | | 162 | | |
| | AAA | TTC | ATG | TCC | ACA | TCC | GTA | GGA | GAC | AGG | GTC | AGC | ATC | ACC | TGC | AAG | GCC | AGT | | | |
| | K | F | M | S | T | S | V | G | D | R | V | S | I | T | C | K | A | S | | | |
| | 171 | | | 180 | | | 189 | | | FR2 | | | 198 | | | 207 | | | 216 | | |
| | CAG | AAT | GTG | ATT | ACT | GCT | GTA | GCC | TGG | TAT | CAA | CAG | AAA | CCA | GGA | CAA | TCT | CCT | | | |
| | Q | N | V | I | T | A | V | A | W | Y | Q | Q | K | P | G | Q | S | P | | | |
| | 225 | | | 234 | | | CDR2 | | | 243 | | | 252 | | | FR3 | | | 261 | | |
| | AAA | TTG | CTG | ATT | TAC | TCG | GCA | TCC | AAT | CGG | TAC | ACT | GGA | GTC | CCT | GAT | CGC | TTC | | | |
| | K | L | L | I | Y | S | A | S | N | R | Y | T | G | V | P | D | R | F | | | |
| | 279 | | | 288 | | | 297 | | | 306 | | | 315 | | | 324 | | | | | |
| | TCA | GGC | AGT | GGG | TCT | GGG | ACA | GAT | TTC | ACT | CTC | ACC | ATC | AGC | AAT | ATG | CAG | TCT | | | |
| | S | G | S | G | S | G | T | D | F | T | L | T | I | S | N | M | Q | S | | | |
| | 333 | | | 342 | | | 351 | | | CDR3 | | | 360 | | | 369 | | | 378 | | |
| | GAA | GAC | CTG | GCA | GAT | TAT | TTC | TGC | CAG | CAA | TAT | AAC | AGC | TAT | CCG | TAC | ACG | TTC | | | |
| | E | D | L | A | D | Y | F | C | Q | Q | Y | N | S | Y | P | Y | T | F | | | |
| | FR4 | | | 387 | | | 396 | | | 405 | | | Bsi WI | | | | | | | | |
| | GGA | GGG | GGG | ACC | AAG | CTG | GAA | ATC | AAA | CGT | ACG | 3' | | | | | | | | | |
| | G | G | G | T | K | L | E | I | K | R | T | | | | | | | | | | |

FIGURE 7

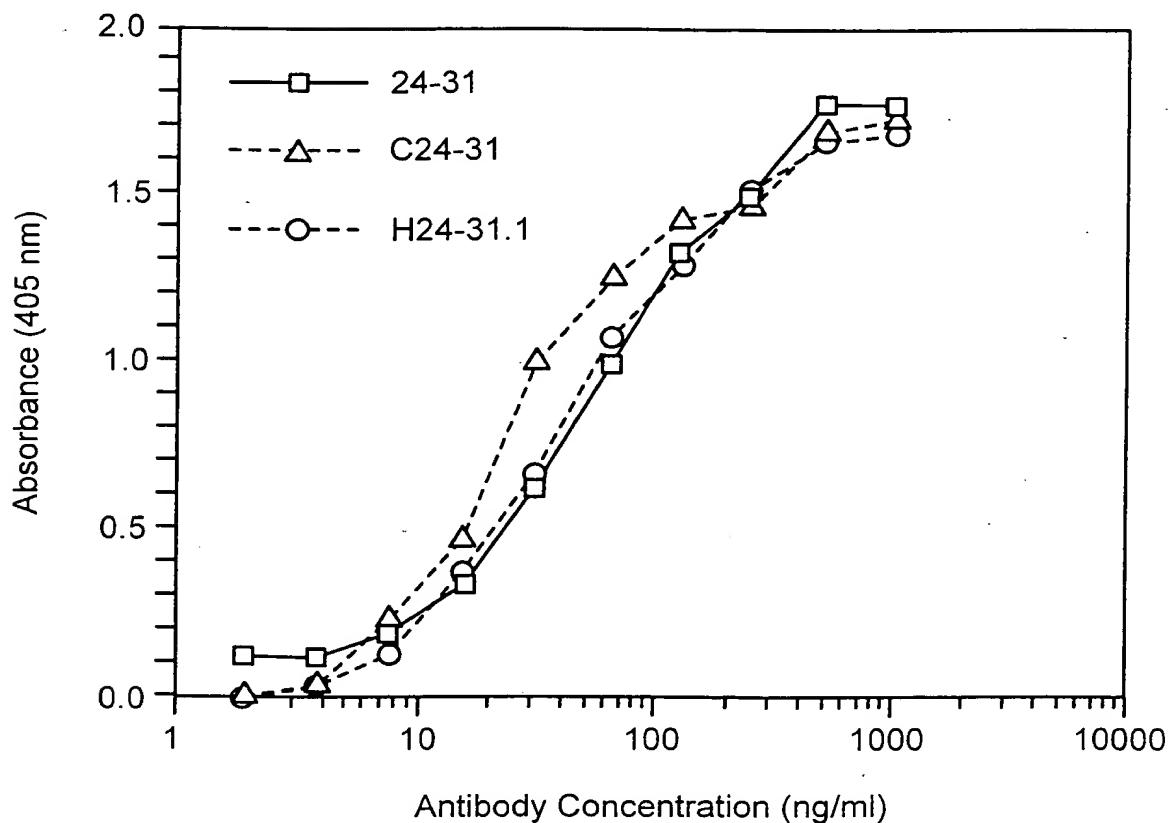
Anti gp39 24-31 V_H Sequence

| | | | | | | | | | | | | | | | | | | |
|----|------|-----|-----|-----|-----|-----|-----|-----|-----|------|-----|-----|-----|-----|-----|------|-----|-----|
| 5' | SalI | | 9 | | 18 | | 27 | | 36 | | 45 | | 54 | | | | | |
| | GTC | GAC | ATG | ATG | GTG | TTA | AGT | CTT | CTG | TAC | CTG | TTG | ACA | GCC | CTT | CCG | GGT | TTC |
| | | | M | M | V | L | S | L | L | Y | L | L | T | A | L | P | G | F |
| | | | +1 | | | | | | | | | | | | | | | |
| | | | 63 | | FRI | 72 | | | 81 | | | 90 | | | 99 | | | 108 |
| | CTG | TCA | GAG | GTG | CAG | CTT | CAG | GAG | TCA | GGA | CCT | AGC | CTC | GTG | AAA | CCT | TCT | CAG |
| | L | S | E | V | Q | L | Q | E | S | G | P | S | L | V | K | P | S | Q |
| | | | 117 | | | 126 | | | 135 | | | 144 | | | 153 | CDR1 | | 162 |
| | ACT | CTG | TCC | CTC | ACC | TGT | TCT | GTC | ACT | GGC | GAC | TCC | ATC | ACT | AAT | GGT | TTC | TGG |
| | T | L | S | L | T | C | S | V | T | G | D | S | I | T | N | G | F | W |
| | | | 171 | FR2 | | 180 | | | 189 | | | 198 | | | 207 | CDR2 | | 216 |
| | ATC | TGG | ATC | CGG | AAA | TTC | CCA | GGG | AAT | AAA | CTT | GAG | TAC | ATG | GGC | TAC | ATA | AGT |
| | I | W | I | R | K | F | P | G | N | K | L | E | Y | M | G | Y | I | S |
| | | | 225 | | | 234 | | | 243 | | | 252 | | | 261 | FR3 | | 270 |
| | TAC | AGT | GGT | AGC | ACT | TAC | TAC | AAT | CCA | TCT | CTC | AAG | AGT | CGA | ATC | TCC | ATC | ACT |
| | Y | S | G | S | T | Y | Y | N | P | S | L | K | S | R | I | S | I | T |
| | | | 279 | | | 288 | | | 297 | | | 306 | | | 315 | | | 324 |
| | CGC | GAC | ACA | TCC | CAG | AAC | CAG | TTC | TAC | CTA | CAA | TTG | AAT | TCT | GTG | ACT | ACT | GAG |
| | R | D | T | S | Q | N | Q | F | Y | L | Q | L | N | S | V | T | T | E |
| | | | 333 | | | 342 | | | 351 | CDR3 | | 360 | | | 369 | | | 378 |
| | GAC | ACA | GGC | ACA | TAT | TAC | TGT | GCC | TGC | CGC | AGT | TAC | GGG | AGG | ACC | CCG | TAC | TAC |
| | D | T | G | T | Y | Y | C | A | C | R | S | Y | G | R | T | P | Y | Y |
| | | | 387 | FR4 | | 396 | | | 405 | | | 414 | | | 423 | NheI | | |
| | TTT | GAC | TTC | TGG | GGC | CAA | GGC | ACC | ACT | CTC | ACC | GTC | TCC | TCA | GCT | AGC | 3' | |
| | F | D | F | W | G | Q | G | T | T | L | T | V | S | S | A | S | | |

FIGURE 8

FIG. 9

Direct Binding of Anti-gp39 Antibodies to mgp39 CHO Cells

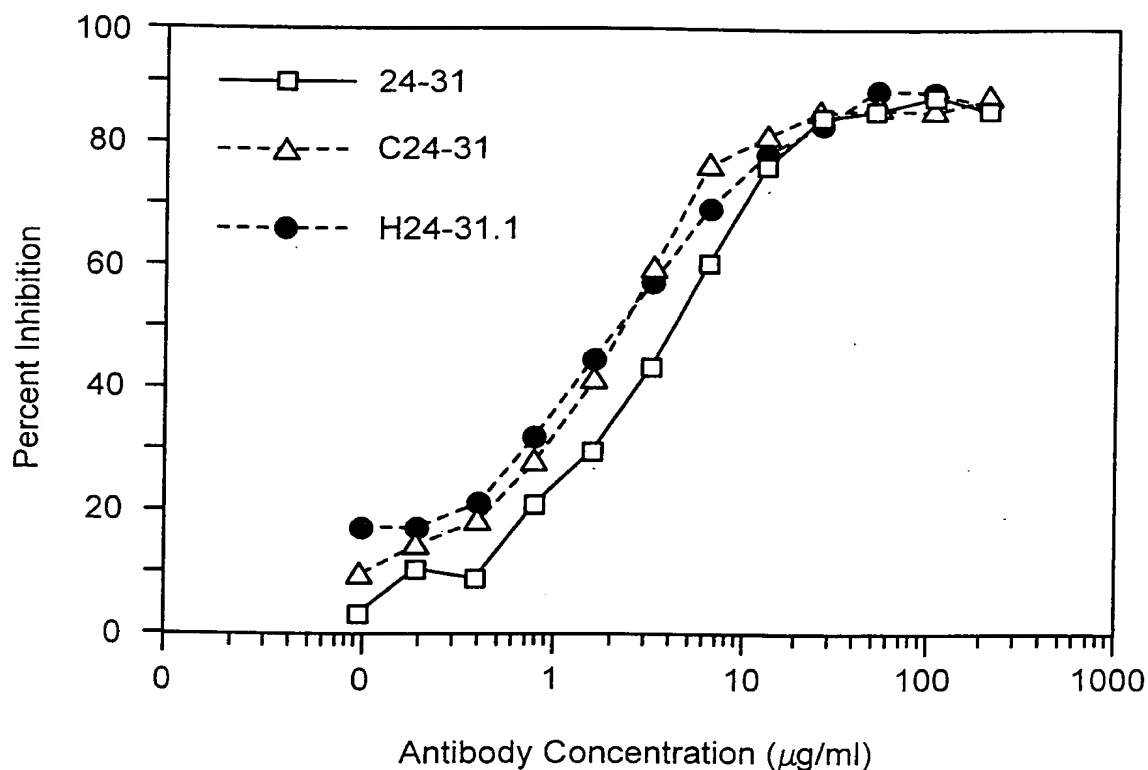


Fifty μ l of 1 μ g/ml solution of each anti-gp39 antibody (murine, chimeric and humanized version 1 of 24-31) was added to wells containing poly-l-lysine fixed mgp39 CHO cells. After a 2 hour incubation, the bound antibodies were detected with either goat anti-human IgG HRP or goat anti-mouse IgG HRP. The binding capacity of each antibody was compared on a plot of absorbance vs antibody concentration.

The figure shows that the half maximal binding in ELISA is achieved at similar concentrations for all three versions, at approximately 40 ng/ml.

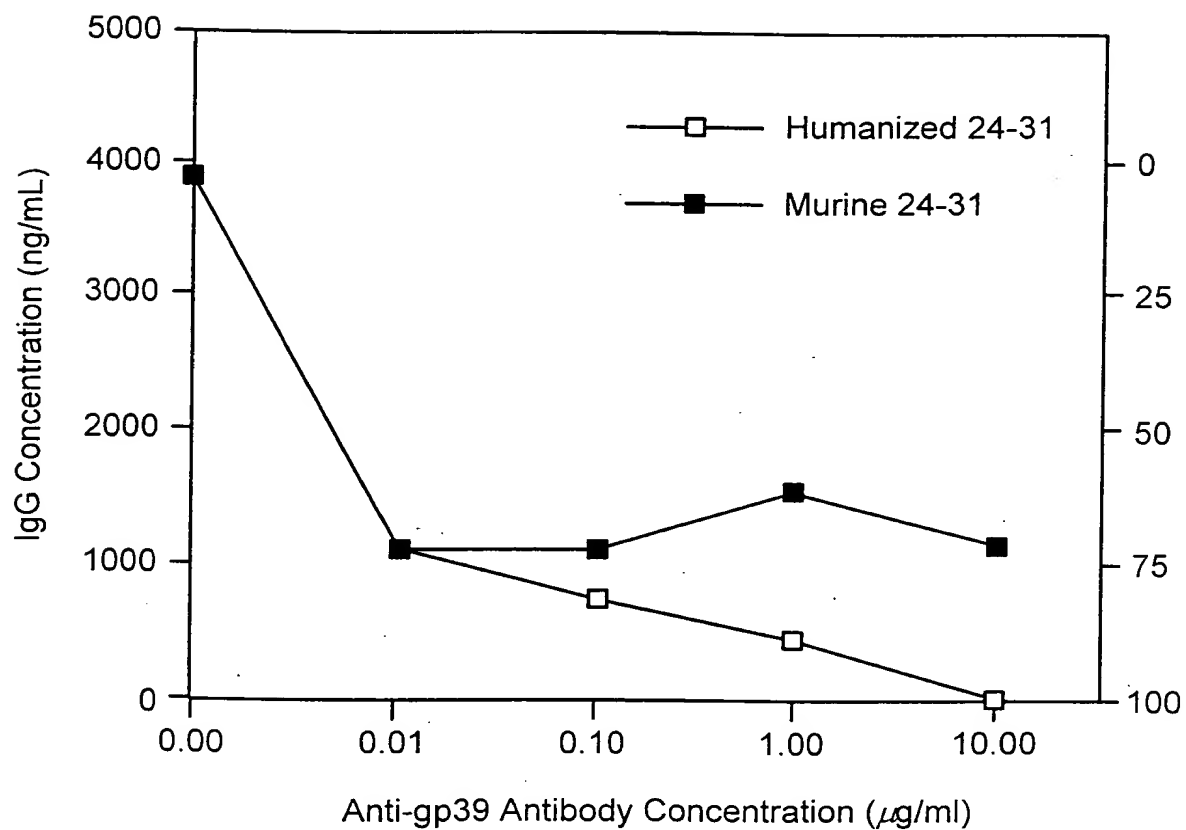
FIG. 10

**Competition Binding of 200 ng/ml Mouse Anti-gp39 Biotin
with Anti-gp39 Antibodies on mgp39 CHO Cells**



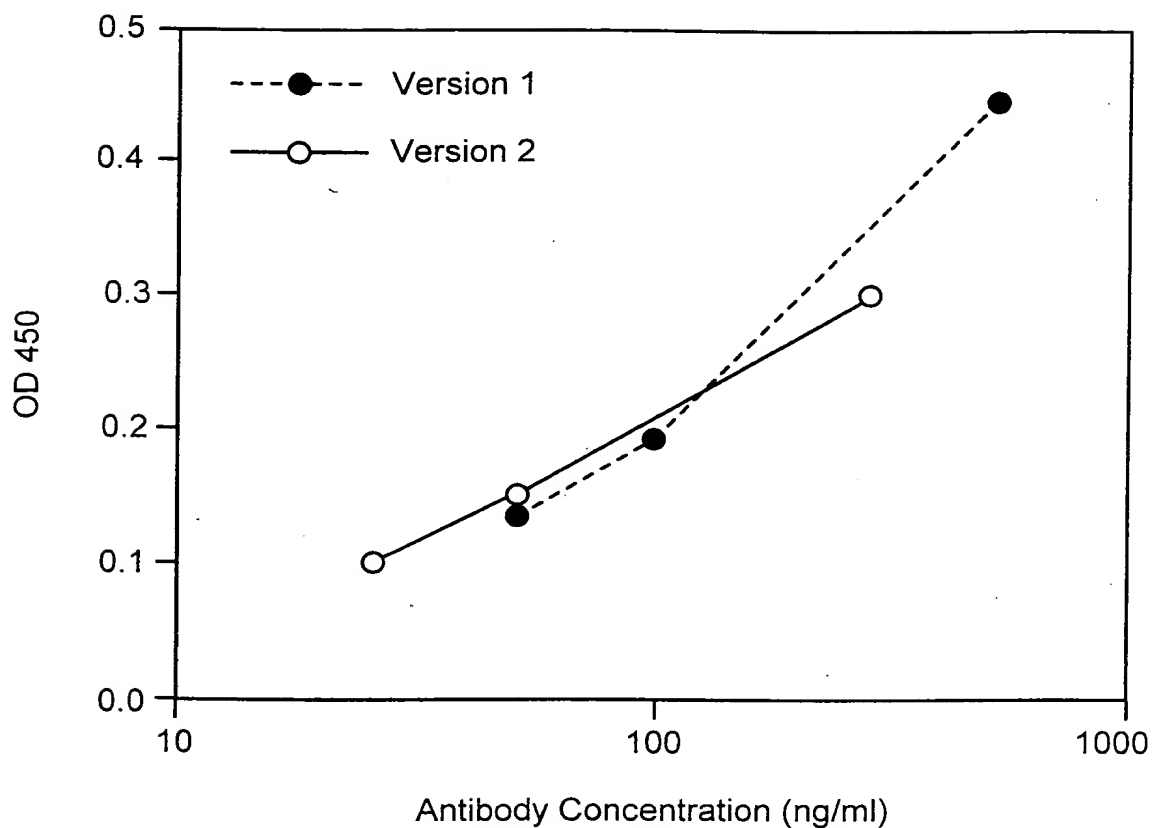
The figure shows that competition for binding to gp39 between biotin labeled 24-31 and the mouse, chimeric and humanized versions 1 are comparable, possibly with the humanized version slightly better than the original antibody, with half-maximal competition at 2 and 4 $\mu\text{g/ml}$, respectively.

FIG. 11



Purified, mitomycin C treated T cells were added into cell culture plates coated with anti-CD3 antibody. Autologous purified B cells were mixed with antibody at described concentrations and added to these plates in regular growth media. After 10 days the supernatant was tested for content of human IgM.

FIG. 12



CHO cell supernatant containing humanized 24-31 version 1 and version 2 in unknown amounts, was incubated on mgp39-CHO cells for 2 hours. After a wash, the amount of bound antibody was determined. The same supernatants were tested in parallel on an ELISA plate coated with Goat α Human γ , to determine the concentration of human IgG present relative to a control of known concentration. The binding data were normalized relative to the total antibody concentration.

FIG. 13

Scatchard Analysis

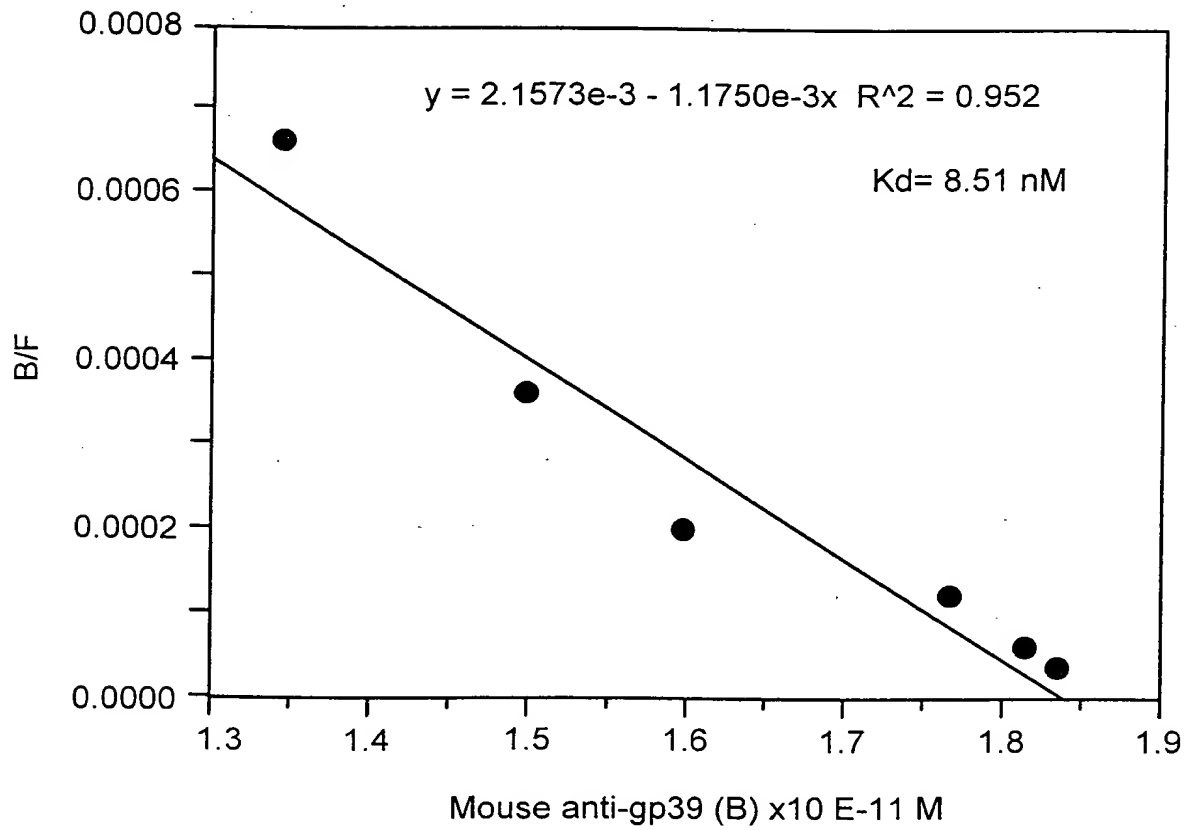


FIG. 14

Scatchard Analysis

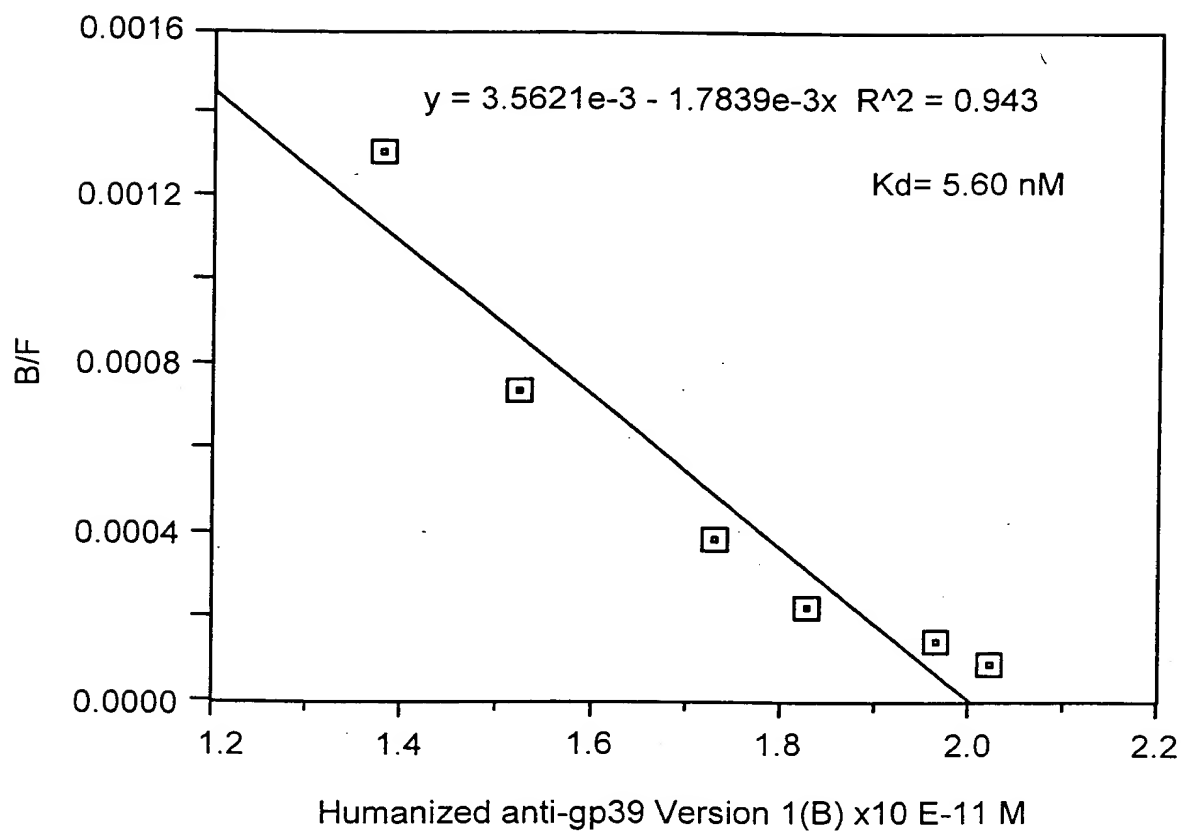
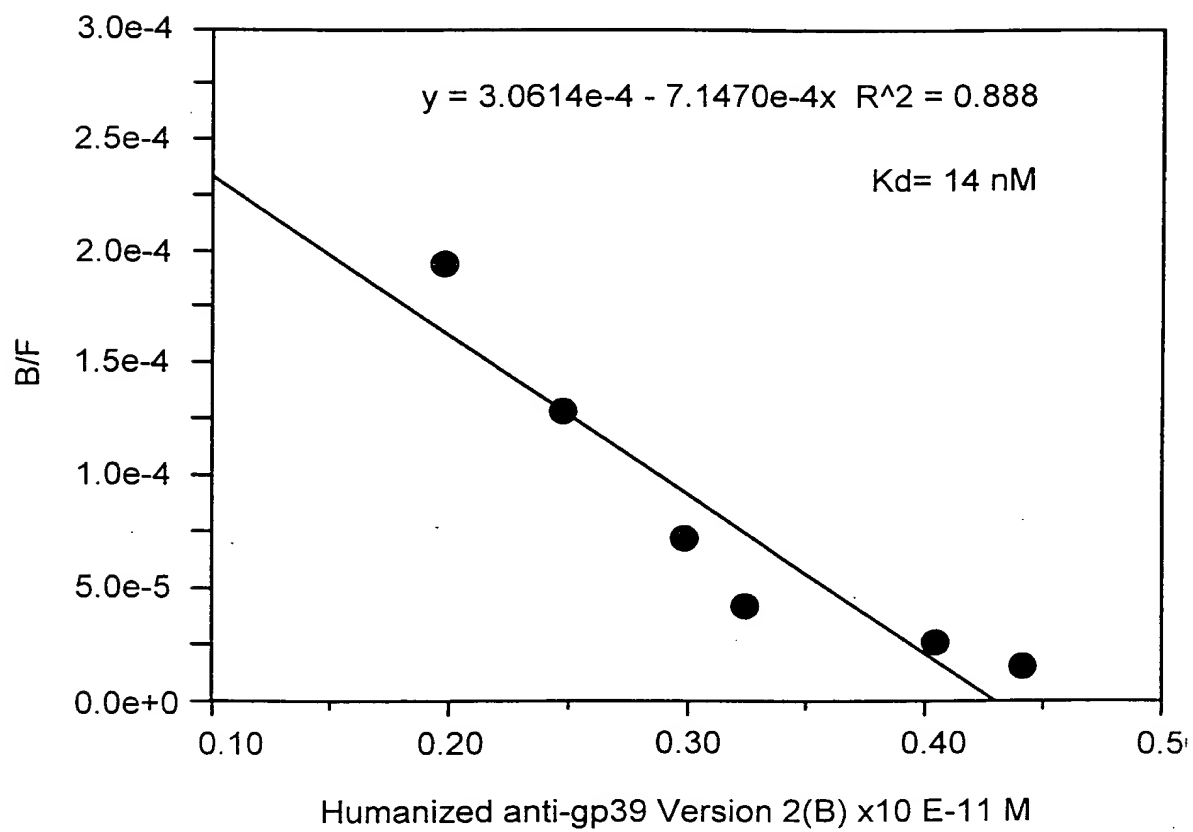


FIG. 15

Scatchard Analysis



Soluble Anti-CD40L Antibody (TRAP1) Stimulates IL-2 Cytokine Release from CD4+ T Cells in Presence of Immobilized Anti-CD3

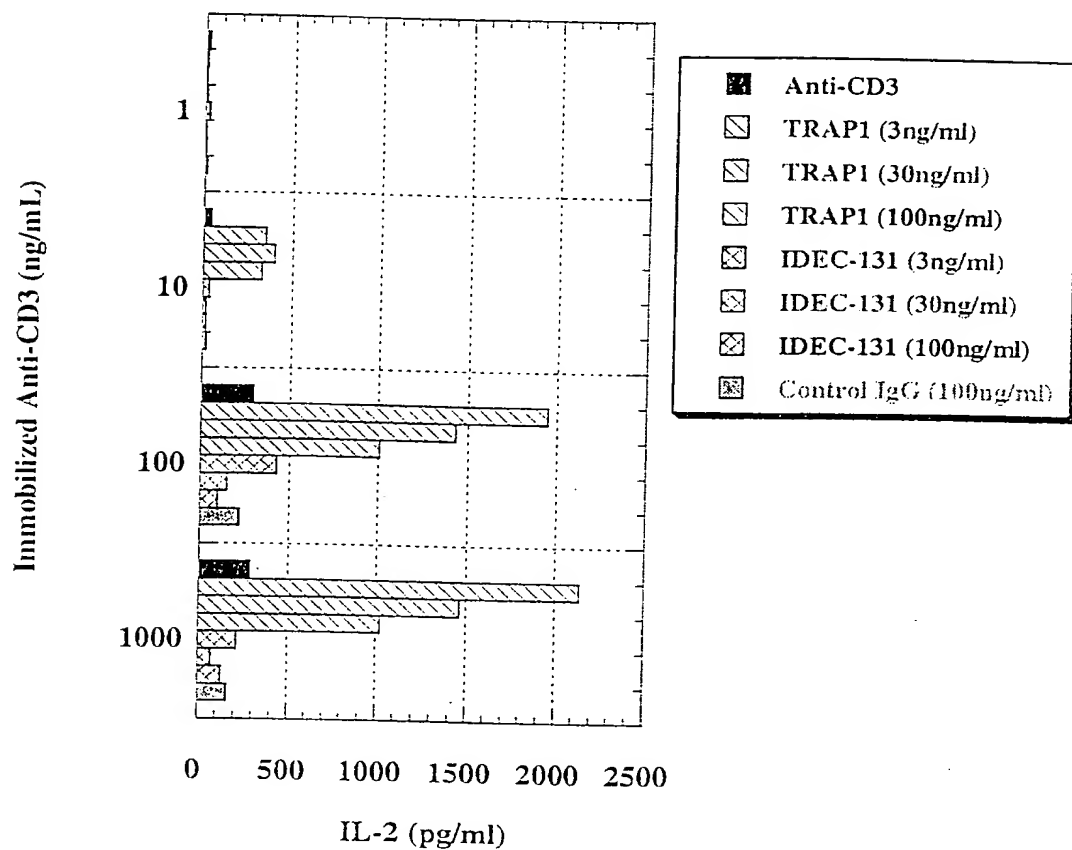


Figure 16

Soluble Anti-CD40L Antibody (TRAP1) Stimulates IL-4 Cytokine Release from CD4+ T cells in Presence of Immobilized Anti-CD3

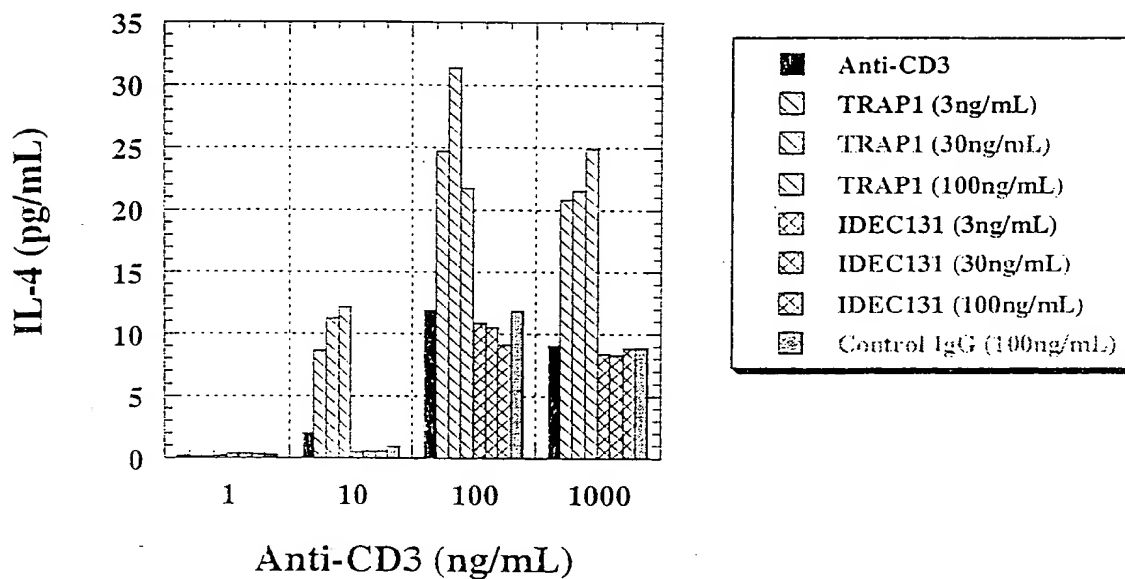


Figure 17

109090-1444250

Stimulation of IL-2 by Agonist TRAP1 Anti-CD40L is Dependent on Co-stimulation with Anti-CD3 and Signaling Through CD40L

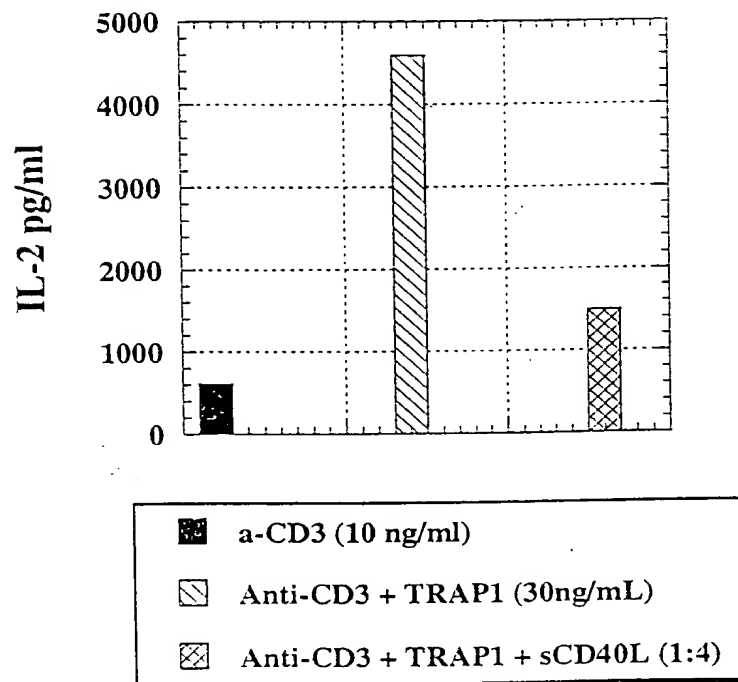


Figure 18

T09090" T4T42860

Soluble Anti-CD40L Antibody (TRAP1) Stimulates Gamma Interferon Release from CD4+ T cells in Presence of Immobilized Anti-CD3

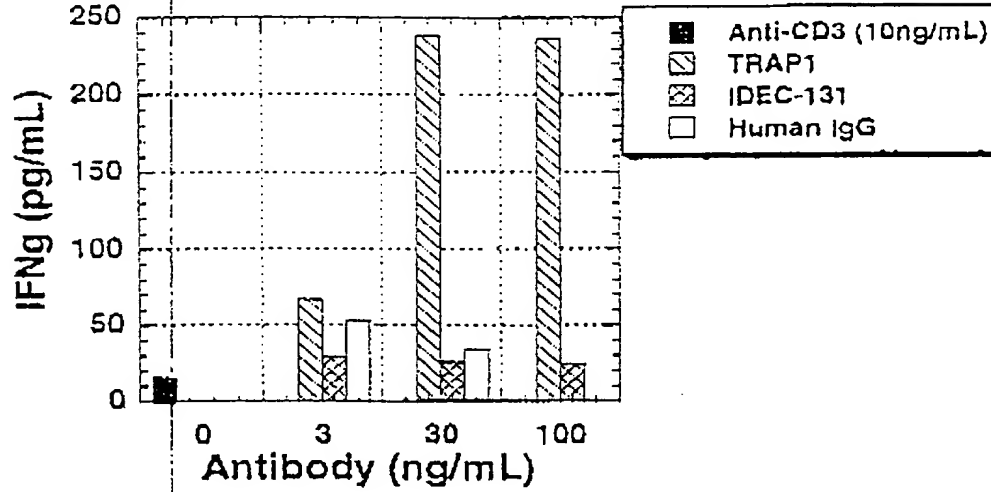


Figure 19

Effect of Anti-CD40L Monoclonal Antibodies on Proliferation of Human CD4+ T cells in vitro

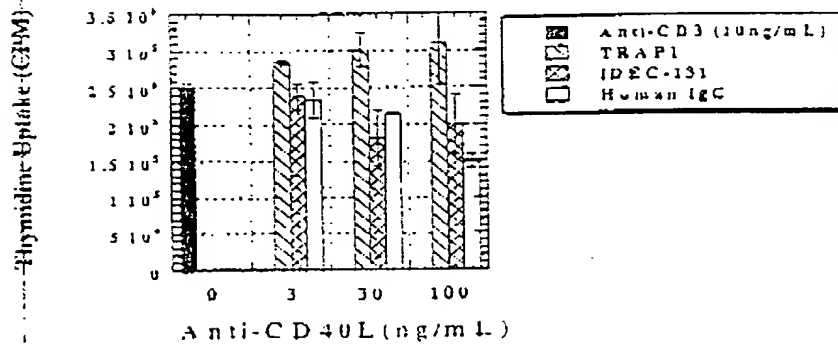


Figure 20